

=====

Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).

Reviewer: Durreshwar Anjum

Timestamp: [year=2008; month=10; day=29; hr=14; min=12; sec=15; ms=896;
]

=====

Reviewer Comments:

<110> APPLICANT: Hakoto KOBAYASHI

Yugo HABATA

Ryo FUJII

Shuji HINUMA

<120> TITLE OF INVENTION: Methods of Screening for Ligands for FPRL2

<130> FILE REFERENCE: 3171 US0P

<140> CURRENT APPLICATION NUMBER:10554234

<141> CURRENT FILING DATE:2005-10-21

<150> PRIOR APPLICATION NUMBER: PCT/JP2004/005829

<151> PRIOR FILING DATE: 2004-04-22

<150> PRIOR APPLICATION NUMBER: JP 2003-118760

<151> PRIOR FILING DATE: 2003-04-23

<160> NUMBER OF SEQ ID NOS: 8

<210> SEQ ID NO 1

<211> LENGTH: 353

<212> TYPE: PRT

<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 1

Please do not insert alpha numeric headings. Please make necessary
changes. This type of error is seen globally throughout the sequence
listing.

Application No: 10554234 Version No: 2.0

Input Set:

Output Set:

Started: 2008-09-30 12:20:13.960
Finished: 2008-09-30 12:20:14.257
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 297 ms
Total Warnings: 3
Total Errors: 0
No. of SeqIDs Defined: 8
Actual SeqID Count: 8

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
W 402	Undefined organism found in <213> in SEQ ID (4)
W 333	tabs used in amino acid numbering SEQID (8)

<110> APPLICANT: Hakoto KOBAYAHSI

Yugo HABATA

Ryo FUJII

Shuji HINUMA

<120> TITLE OF INVENTION: Methods of Screening for Ligands for FPRL2

<130> FILE REFERENCE: 3171 USOP

<140> CURRENT APPLICATION NUMBER:10554234

<141> CURRENT FILING DATE:2005-10-21

<150> PRIOR APPLICATION NUMBER: PCT/JP2004/005829

<151> PRIOR FILING DATE: 2004-04-22

<150> PRIOR APPLICATION NUMBER: JP 2003-118760

<151> PRIOR FILING DATE: 2003-04-23

<160> NUMBER OF SEQ ID NOS: 8

<210> SEQ ID NO 1

<211> LENGTH: 353

<212> TYPE: PRT

<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 1

```
Met Glu Thr Asn Phe Ser Ile Pro Leu Asn Glu Thr Glu Glu Val Leu
      5              10              15
Pro Glu Pro Ala Gly His Thr Val Leu Trp Ile Phe Ser Leu Leu Val
      20              25              30
His Gly Val Thr Phe Val Phe Gly Val Leu Gly Asn Gly Leu Val Ile
      35              40              45
Trp Val Ala Gly Phe Arg Met Thr Arg Thr Val Asn Thr Ile Cys Tyr
      50              55              60
Leu Asn Leu Ala Leu Ala Asp Phe Ser Phe Ser Ala Ile Leu Pro Phe
      65              70              75              80
Arg Met Val Ser Val Ala Met Arg Glu Lys Trp Pro Phe Ala Ser Phe
      85              90              95
Leu Cys Lys Leu Val His Val Met Ile Asp Ile Asn Leu Phe Val Ser
      100             105             110
Val Tyr Leu Ile Thr Ile Ile Ala Leu Asp Arg Cys Ile Cys Val Leu
      115             120             125
His Pro Ala Trp Ala Gln Asn His Arg Thr Met Ser Leu Ala Lys Arg
      130             135             140
Val Met Thr Gly Leu Trp Ile Phe Thr Ile Val Leu Thr Leu Pro Asn
      145             150             155             160
Phe Ile Phe Trp Thr Thr Ile Ser Thr Thr Asn Gly Asp Thr Tyr Cys
      165             170             175
Ile Phe Asn Phe Ala Phe Trp Gly Asp Thr Ala Val Glu Arg Leu Asn
      180             185             190
Val Phe Ile Thr Met Ala Lys Val Phe Leu Ile Leu His Phe Ile Ile
      195             200             205
Gly Phe Thr Val Pro Met Ser Ile Ile Thr Val Cys Tyr Gly Ile Ile
      210             215             220
Ala Ala Lys Ile His Arg Asn His Met Ile Lys Ser Ser Arg Pro Leu
      225             230             235             240
Arg Val Phe Ala Ala Val Val Ala Ser Phe Phe Ile Cys Trp Phe Pro
      245             250             255
Tyr Glu Leu Ile Gly Ile Leu Met Ala Val Trp Leu Lys Glu Met Leu
      260             265             270
Leu Asn Gly Lys Tyr Lys Ile Ile Leu Val Leu Ile Asn Pro Thr Ser
      275             280             285
Ser Leu Ala Phe Phe Asn Ser Cys Leu Asn Pro Ile Leu Tyr Val Phe
```

290	295	300
Met Gly Arg Asn Phe Gln Glu Arg Leu Ile Arg Ser Leu Pro Thr Ser		
305	310	315
Leu Glu Arg Ala Leu Thr Glu Val Pro Asp Ser Ala Gln Thr Ser Asn		320
	325	330
Thr His Thr Thr Ser Ala Ser Pro Pro Glu Glu Thr Glu Leu Gln Ala		335
	340	345
		350
Met		

<210> SEQ ID NO 2

<211> LENGTH: 1059

<212> TYPE: DNA

<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 2

atggaaacca	acttctccat	tcctctgaat	gaaactgagg	aggtgctccc	tgagcctgct	60
ggccacaccg	ttctgtggat	cttctcattg	ctagtccacg	gagtcacctt	tgtcttcggg	120
gtcctgggca	atgggcttgt	gatctgggtg	gctggattcc	ggatgacacg	cacagtcaac	180
accatctggt	acctgaacct	ggccctagct	gacttctctt	tcagtgccat	cctaccattc	240
cgaatggtct	cagtcgccat	gagagaaaaa	tggccttttg	cgtcattcct	atgtaagtta	300
gttcatgtta	tgatagacat	caacctgttt	gtcagtgtct	acctgatcac	catcattgct	360
ctggaccgct	gtatttgtgt	cctgcatcca	gcctgggccc	agaaccatcg	caccatgagt	420
ctggccaaga	gggtgatgac	gggactctgg	attttcacca	tagtccttac	cttaccaaat	480
ttcatcttct	ggactacaat	aagtactacg	aatggggaca	catactgtat	tttcaacttt	540
gcattctggg	gtgacactgc	tgtagagagg	ttgaacgtgt	tcattaccat	ggccaaggtc	600
tttctgatcc	tccacttcat	tattggcttc	acggtgccta	tgtccatcat	cacagtctgc	660
tatgggatca	tcgctgccaa	aattcacaga	aaccacatga	ttaaattccag	ccgtccctta	720
cgtgtcttcg	ctgctgtggg	ggcttctttc	ttcatctgtt	ggttccctta	tgaactaatt	780
ggcattctaa	tggcagtcct	gctcaaagag	atggtgttaa	atggcaaata	caaaatcatt	840
cttgtcctga	ttaacccaac	aagctccttg	gcctttttta	acagctgcct	caacccaatt	900
ctctacgtct	ttatgggtcg	taacttccaa	gaaagactga	ttcgctcttt	gcccactagt	960
ttggagaggg	ccctgactga	gggccctgac	tcagcccaga	ccagcaacac	acacaccact	1020
tctgcttcac	ctcctgagga	gacggagtta	caagcaatg			1059

<210> SEQ ID NO 3

<211> LENGTH: 6

<212> TYPE: PRT

<213> ORGANISM: Artificial Sequence

<220> FEATURE:

<223> OTHER INFORMATION: amino acid sequence of GHRP-6

<220> FEATURE:

<223> OTHER INFORMATION: Trp is a D-form

<400> SEQUENCE: 3

His Trp Ala Trp Phe Lys
1 5

<210> SEQ ID NO 4

<211> LENGTH: 11

<212> TYPE: PRT

<213> ORGANISM: Aplysia sp.

<400> SEQUENCE: 4

Ala Arg Pro Gly Tyr Leu Ala Phe Pro Arg Met
1 5 10

<210> SEQ ID NO 5

<211> LENGTH: 12

<212> TYPE: PRT
<213> ORGANISM: Sus scrofa
<400> SEQUENCE: 5
Met Pro His Ser Phe Ala Asn Leu Pro Leu Arg Phe
1 5 10

<210> SEQ ID NO 6
<211> LENGTH: 36
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens
<400> SEQUENCE: 6
Tyr Pro Ser Lys Pro Asp Asn Pro Gly Glu Asp Ala Pro Ala Glu Asp
1 5 10 15
Leu Ala Arg Tyr Tyr Ser Ala Leu Arg His Tyr Ile Asn Leu Ile Thr
20 25 30
Arg Gln Arg Tyr
35

<210> SEQ ID NO 7
<211> LENGTH: 10
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens
<400> SEQUENCE: 7
Gly Asn His Trp Ala Val Gly His Leu Met
1 5 10

<210> SEQ ID NO 8
<211> LENGTH: 6
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens
<400> SEQUENCE: 8
Met Val Met Tyr Lys Trp
1 5